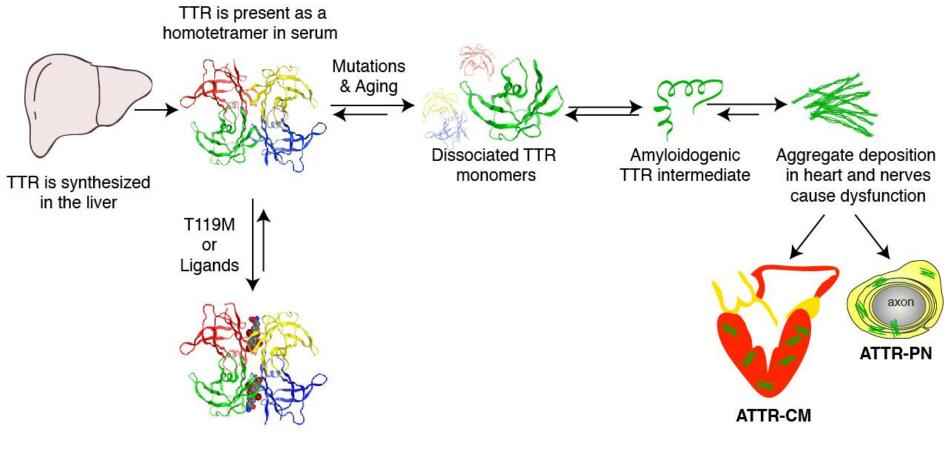


AG10 Stabilizes Pathogenic TTR Variants With High Potency – Potential for an Effective Treatment for ATTR Cardiomyopathy

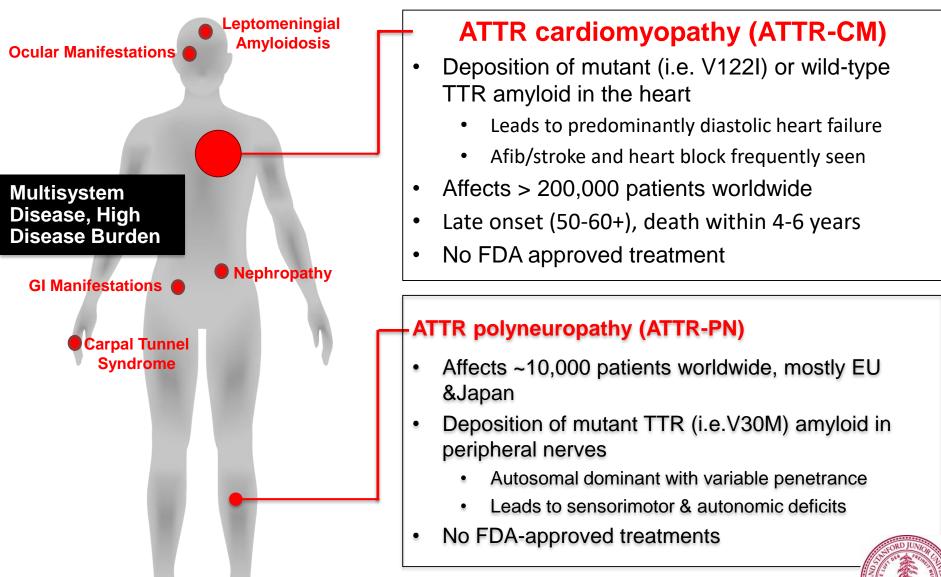


ATTR-Cardiomyopathy (CM) And ATTR-Polyneuropathy (PN) Are Caused By Aggregation Of Misfolded TTR Monomers



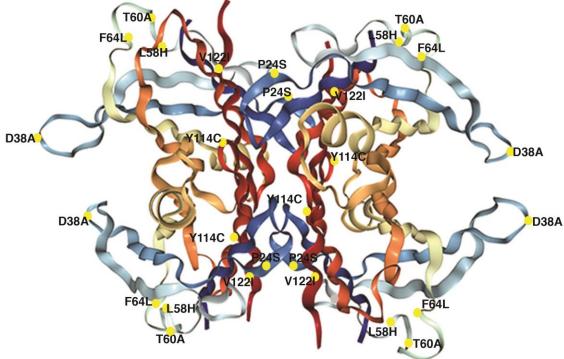


TTR amyloidosis is a systemic disorder



Source: Grogan, M et al. *JACC* 2016, 68:1014-20; Planté-Bordeneuve, V. et al, *Lancet Neurol* 2011, 10:1086-97

Does AG10 Stabilize a Broad Range of Pathogenic TTR Variants?



V122I: Cardiomyopathy

T60A: Cardiomyopathy & polyneuropathy

P24S: Cardiomyopathy & polyneuropathy

D38A: Cardiomyopathy & polyneuropathy

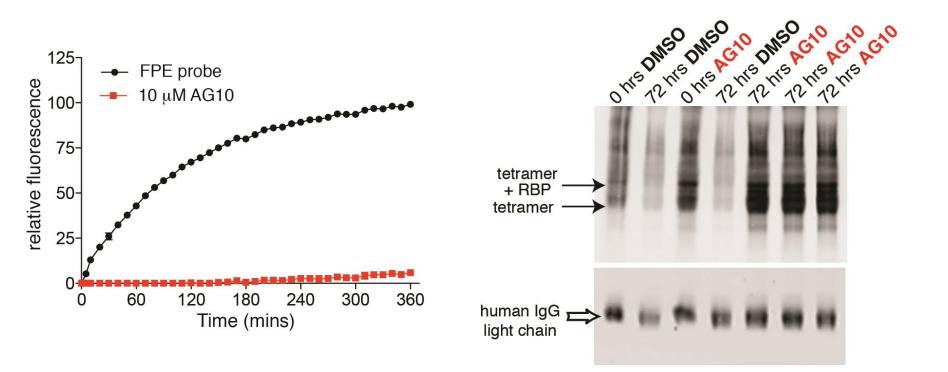
L58H:Cardiomyopathy & polyneuropathy

F64L:Polyneuropathy

Y114C: Polyneuropathy with leptomeningeal complications

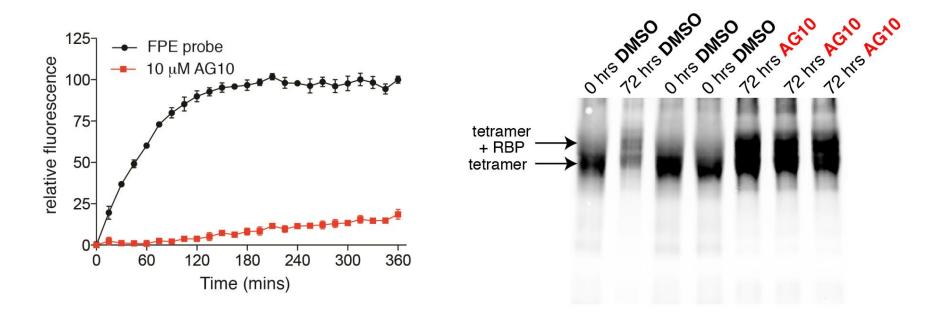


AG10 Stabilizes Mutant TTR from V122I ATTR Patients



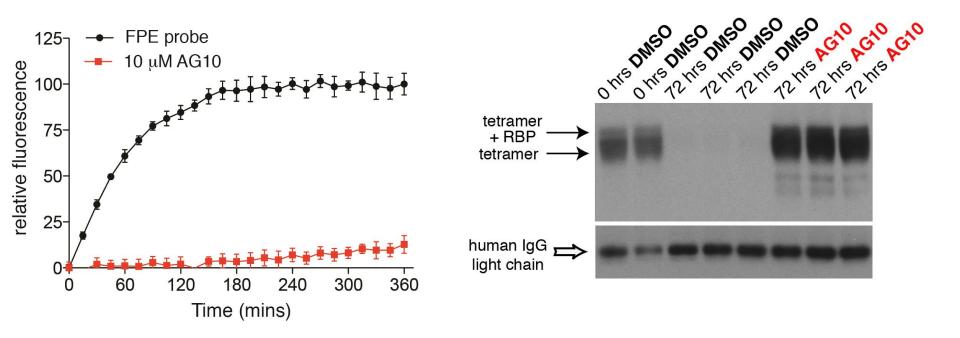
FPE Assay: Change in fluorescence due to modification of TTR in human serum by a covalent probe, which becomes fluorescent following binding to TTR. The lower the binding of the probe/fluorescence the higher the binding selectivity and affinity of the ligand to TTR. AG10 was used at 10 μ M

AG10 Stabilizes Mutant TTR from T60A ATTR Patients



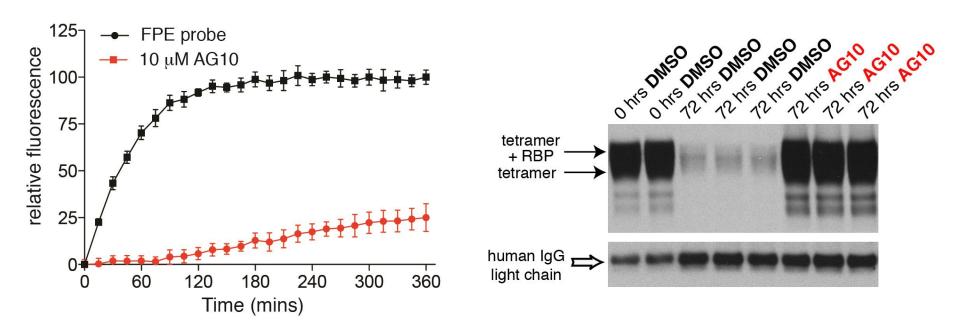


AG10 Stabilizes Mutant TTR from P24S ATTR Patients



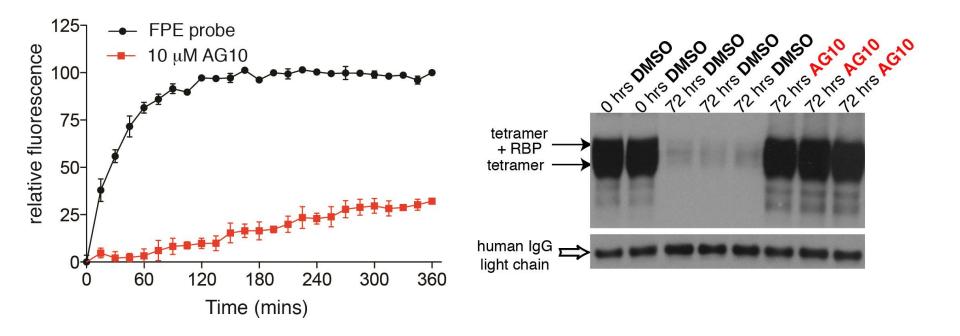


AG10 Stabilizes Mutant TTR from D38A ATTR Patients



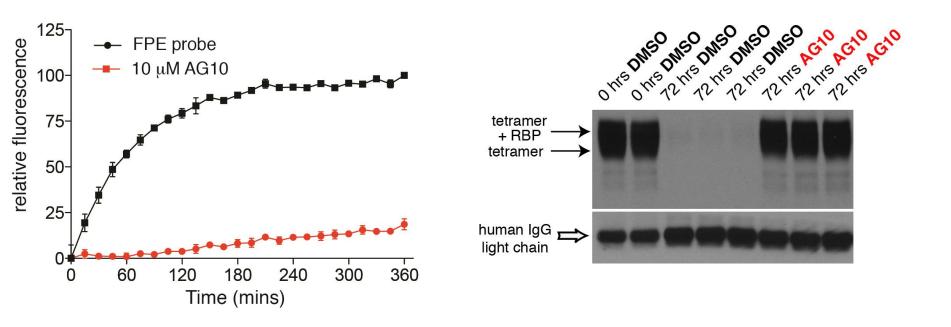


AG10 Stabilizes Mutant TTR from L58H ATTR Patients



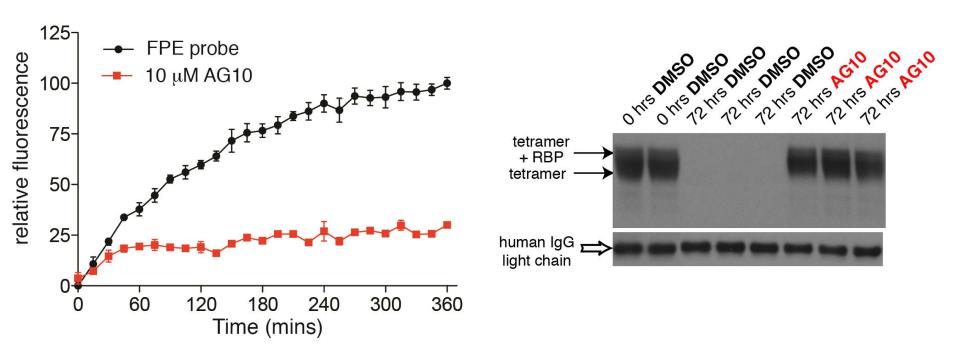


AG10 Stabilizes Mutant TTR from F64L ATTR Patients



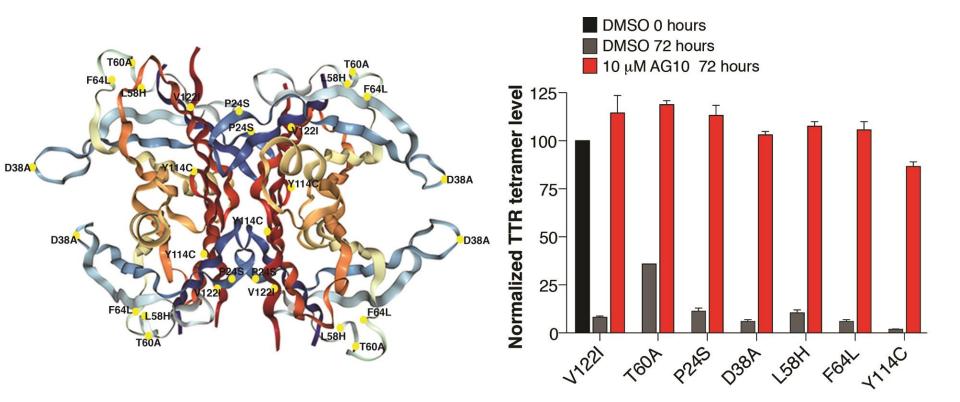


AG10 Stabilizes Mutant TTR from Y114C ATTR Patients





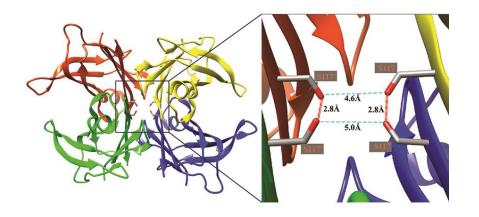
Does AG10 Stabilize a Broad Range of Pathogenic TTR Variants? - YES





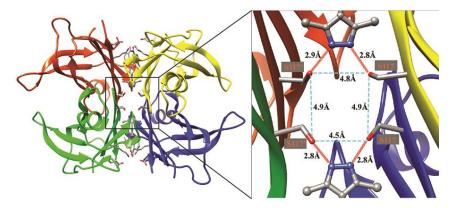
AG10 has a unique binding mode, which mimics the effect of the TTR trans-suppressor mutation - T119M

The naturally occurring trans-suppressor mutation T119M super-stabilizes TTR AG10 binding to TTR mimics the stabilizing interactions of T119M variant to S117



- The T119M polymorphism creates H-bonds within the complex that super-stabilize the TTR tetramer and functions as a trans-suppressor mutation in V30M carriers.
- T119M heterozygotes have a 5-10 year longer life-span and significantly lower risk of cerebrovascular disease

Hammarström et al, *Science*, 2001, 293:2459-62 Hornstrup et al, *Arterioscler Thromb Vasc Biol*, 2013, 33(6), 1441-7 Penchala et al. *Proc Natl Acad Sci USA*, 2013, 110:9992-7



- AG10 mimics the structural effects of T119M.
- Stabilization of TTR by AG10 may mimic the clinical effect and lead to improved outcomes

Sebastiao et al, *J. Mol.Biol.* 2001, 306, 733-44 Miller et al, unpublished data



Coworkers and Collaborators

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Stanford University:

Ron Witteles Michaela Liedtke **University of the Pacific:**

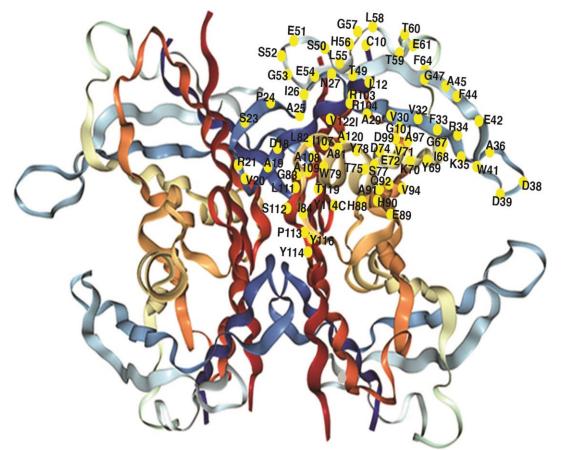
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Mark Miller



TTR Mutations

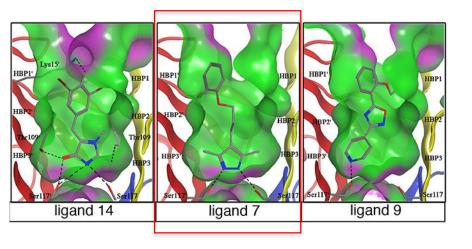


- > 100 mutations in the TTR gene have been found to cause TTR amyloidosis (ATTR).
- Most of these alter the TTR structure, resulting in either kinetic or thermodynamic destabilization
- The most common ATTR mutations are V122I (3.4% of African Americans) and V30M.

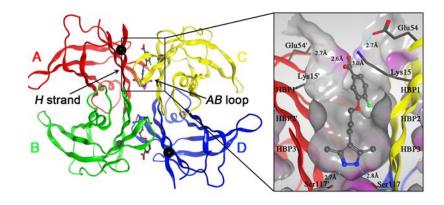


Discovery and Development of AG10

- HTS of 130,000 identified 32 compounds with IC50 <1 μM
- Crystal structure of top novel ligands



 The crystal structure of Ligand 7 was used as a starting point for SAR studies. • AG10 was the most potent analogue with the best physicochemical properties

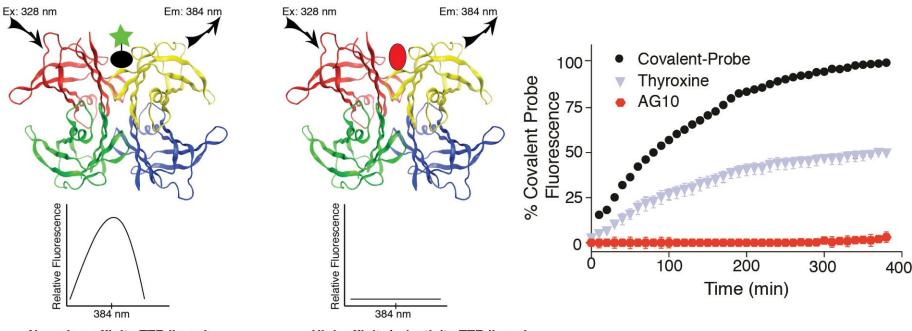


- AG10 was selected for ADME and Toxicity studies
- IND was filed in August 2017 and Phase
 1 clinical studies started in September
 2017



Alhamadsheh et al, *Science Transl Med*, 2011, August 24, Vol.97 Penchala S. et al. *Proc Natl Acad Sci USA*, 2013, 110:9992-7

AG10 binds with high affinity and high selectivity to human serum TTR



No or low affinity TTR ligand: covalent probe forms an amide bond with Lys-15 of TTR, creating a fluorescent conjugate. **High affinity/selectivity TTR ligand** competes effectively with the FPE probe and prevents the formation of a fluorescent conjugate.

